

IN THE CLAIMS

Please replace Claim 13 with the following rewritten claim:

- 13. (Amended) A method according to claim 1, wherein said RPP has a sequence selected from the group consisting of the sequences depicted in Figure 3 (SEQ ID NOS:1-6). –

Please replace Claim 14 with the following rewritten claim:

- 14. (Amended) A method according to claim 1, wherein said RPP has a sequence selected from the group consisting of the sequences depicted in Figure 4 (SEQ ID NOS:7-13).–

Please replace Claim 24 with the following rewritten claim:

- 24. (Amended) A method of quantifying the amount of a plurality of germline constructs comprising:
  - a) preparing mRNA from said plurality of cells to form an mRNA mixture;
  - c) adding at least three RNase protection probes (RPPs) selected from the group consisting of the sequences depicted in Figures 3 and 4 (SEQ ID NOS:1-13);
  - d) adding an RNase protection enzyme (RPE) to said mixture, such that mRNA that is not protected is digested;
  - e) quantifying the amount of said germline mRNA.–

Please replace Claim 25 with the following rewritten claim:

- 25. (Amended) A kit for quantifying the amount of germline mRNA in a sample, comprising
  - a) at least one RNase protection probe (RPP) comprising a nucleic acid sequence selected from the group consisting of the nucleic acid sequences of the Ig $\alpha$ 1, Ig $\alpha$ 2, Ig-epsilon, Ig gamma-1, Ig gamma-2, Ig gamma-3 and Ig gamma-4 RPPs set forth in Figures 3 and 4 (SEQ ID NOS:1-13); and
  - b) an RNase protection enzyme (RPE);and optionally comprising at least one RNase protection probe (RPP) which is substantially complementary to a transcript of a housekeeping gene.–

REMARKS

The specification and claims have been amended to include a Sequence Listing and proper reference to the sequences therein. Attached hereto is a marked-up version of the changes made to the specification and claims by the current amendment. The attached page is captioned “Version with markings to show changes made.”